

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/817/198A

CRF Processing Date: 7/31/2002
 Edited by: [Signature]
 Verified by: [Signature] (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: globally deleted format markers and inserted hard returns

RECEIVED

AUG 06 2002

TECH CENTER 1600/2900



1600

RAW SEQUENCE LISTING

DATE: 07/31/2002

PATENT APPLICATION: US/09/817,198A

TIME: 18:54:19

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07312002\I817198A.raw

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4 <110> APPLICANT: YE, Jane et al.
6 <120> TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
7   NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
8   PROTEINS, AND USES THEREOF
10 <130> FILE REFERENCE: CL001188
12 <140> CURRENT APPLICATION NUMBER: 09/817,198A
13 <141> CURRENT FILING DATE: 2001-03-27
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92 35 40 45
93 Thr Ile Glu Val Asp Gly Ile Lys Val Arg Ile Gln Ile Trp Asp Thr
94 50 55 60
95 Ala Gly Gln Glu Arg Tyr Gln Thr Ile Thr Lys Gln Tyr Tyr Arg Arg
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100 100 105 110
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102 115 120 125
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243 ctgggggcaa atgcagtgtg acgttgtgag agggtcaggg ctgggtctgt gtcagccttc 7260
244 aggcagcctg agaccagtct ctacctaact tgttcccctg gtacctagaa aggaaggga 7320
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252 cggctattat tattattcct agctataaga atgctgtaga gatgaataca ctgtcagtga 7800
253 gctaggaggt catcctgtgt atccatcact tgtgcactca gtcgttcagg cgctatttgc 7860
254 tgaacaccaa ctacatgcca ggtgccatgc taagatttgg ggacacagtg gtgacaaaa 7920

```

VERIFICATION SUMMARY

DATE: 07/31/2002

PATENT APPLICATION: US/09/817,198A

TIME: 18:54:20

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07312002\I817198A.raw



1600

RAW SEQUENCE LISTING

DATE: 07/31/2002

PATENT APPLICATION: US/09/817,198A

TIME: 18:41:31

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07312002\I817198A.raw

Does Not Comply
Corrected Diskette Needed

partial
listing
of global error

4 <110> APPLICANT: YE, Jane et al.
6 <120> TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
7 NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
8 PROTEINS, AND USES THEREOF
10 <130> FILE REFERENCE: CL001188
12 <140> CURRENT APPLICATION NUMBER: 09/817,198A
13 <141> CURRENT FILING DATE: 2001-03-27
15 <160> NUMBER OF SEQ ID NOS: 33
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

676 <210> SEQ ID NO: 6
677 <211> LENGTH: 4
678 <212> TYPE: PRT
679 <213> ORGANISM: Homo sapien
681 <400> SEQUENCE: 6
E--> 682 Asn Ser Ser Lys 1
684 <210> SEQ ID NO: 7
685 <211> LENGTH: 4
686 <212> TYPE: PRT
687 <213> ORGANISM: Homo sapien
689 <400> SEQUENCE: 7
E--> 690 Thr Asp Asn Glu 1
692 <210> SEQ ID NO: 8
693 <211> LENGTH: 4
694 <212> TYPE: PRT
695 <213> ORGANISM: Homo sapien
697 <400> SEQUENCE: 8
E--> 698 Ser Asp Val Asp 1
700 <210> SEQ ID NO: 9
701 <211> LENGTH: 9
702 <212> TYPE: PRT
703 <213> ORGANISM: Homo sapien
705 <400> SEQUENCE: 9
E--> 706 Lys Trp Val Ser Asp Val Asp Glu Tyr 1
708 <210> SEQ ID NO: 10
709 <211> LENGTH: 6
710 <212> TYPE: PRT
711 <213> ORGANISM: Homo sapien
713 <400> SEQUENCE: 10
E--> 714 Gly Val Gly Lys Thr Cys 1

(global errors)
delete format marker and insert hard return

same error

same

same

same

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/817,198A

DATE: 07/31/2002
TIME: 18:41:32

Input Set : A:\Pto.amc
Output Set: N:\CRF3\07312002\I817198A.raw

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716 <210> SEQ ID NO: 11
717 <211> LENGTH: 6
718 <212> TYPE: PRT
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721 <400> SEQUENCE: 11
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724 <210> SEQ ID NO: 12
725 <211> LENGTH: 8
726 <212> TYPE: PRT
727 <213> ORGANISM: Homo sapien
729 <400> SEQUENCE: 12
E--> 730 Gly Asp Ser Gly Val Gly Lys Thr 1            5
732 <210> SEQ ID NO: 13
733 <211> LENGTH: 14
734 <212> TYPE: PRT
735 <213> ORGANISM: Homo sapien
737 <400> SEQUENCE: 13
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Leu Leu Leu Ile Gly Asp Ser Gly Val Gly Lys Thr Cys Leu 1    5          10
761 <210> SEQ ID NO: 15
762 <211> LENGTH: 601
763 <212> TYPE: DNA
764 <213> ORGANISM: Homo sapien
766 <400> SEQUENCE: 15
767 tattaaggga ctgggattc tcccttatct tgggcgtgtt tttcagcatt aactaaaact      60
768 taaaggaaaag agttggatgg tcaagaaaag ctttttcctt aagtgatatg gacagtttct      120
769 caaggaggta gaagggggcag ccaggagaca aatcaaggag ccaacgaaat gagtgtctacc      180
770 aagtcatagt cattcgctta tttttaaaaa atgcgtgtcc tgtatgccag gctctgcact      240
771 gagaccgaga gattccaaga tgaataatac ctacagtcac tgttctcaaa ttgtgcatta      300
772 yctaaaacac attacatgac catgctggcc actgacgag gcacctttcc caggggcttt      360
773 ttttgtgaat taagaaaaca aggttaattca ccagttattg ccaagatagt ttggcttctt      420
774 ggctcatgtg gatatcacct aggccagtac ttttgtgatt tactgtgtac tccactttaa      480
E--> 775
cggcctgcga ttttctagag aagaacccgc cagggagcag tgagaggcct ccctggtaga      540ctgagacact gactgtccct
777 <210> SEQ ID NO: 16
778 <211> LENGTH: 601
779 <212> TYPE: DNA
780 <213> ORGANISM: Homo sapien
782 <400> SEQUENCE: 16
783 atgccagggtg ccatgctaag atttggggac acagtgggtga caaaaacaga cagaaaccaa      60
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785 ctagtacccc ctttccctg gcagtgccag ggtctgagaa ggaagagtga ggtggtgagg      180
786 aggtgtgaag cagtgggggtg acctgagagg agaggatggg gtggctttgc ctcaaggctt      240
787 gggcccctgc taggtgtcgc tctgcctcag gcctctgttt ctctcctga cacaggcaca      300
788 ractcggcct cccaccctt ccccaaggac atgaccttgg gaaggaacat atctgaagcc      360
789 cgcggagggt ttccgctgct gtgcatctgt gccacagatc cgcagatgca cccacagctg      420
790 ggagcaccgg ttctcccgcc ctacctgcac tccctggttt ctgttccttc ctctcctcc      480
E--> 791
ttccttctcc ccgtcccca gacaggctgg tgatgagctt tataacatga aagctgatat      540ttggccatta tccttctacc
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794 <211> LENGTH: 601
795 <212> TYPE: DNA

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